

OIPE

RAW SEQUENCE LISTING

PATENT APPLICATION: US/09/973,451

DATE: 12/04/2001

TIME: 20:18:50

Input Set : N:\Crf3\RULE60\09973451.raw
 Output Set: N:\CRF3\12042001\I973451.raw

PS

ENTERED

1 <110> APPLICANT: JACOBSON, Myron K.
 2 JACOBSON, Elaine L.
 3 AM, Jean-Christophe
 4 LIN, Winston
 5 <120> TITLE OF INVENTION: GENES ENCODING SEVERAL POLY(ADP-RIBOSE) GLYCOHYDROLASE
 6 (PARG) ENZYMES,
 7 THE PROTEINS AND FRAGMENTS THEREOF, AND ANTIBODIES IMMUNOREACTIVE
 8 THEREWITH
 9 <130> FILE REFERENCE: NIAD 201
 10 <140> CURRENT APPLICATION NUMBER: 09/973,451
 11 <141> CURRENT FILING DATE: 2001-10-09
 12 <150> PRIOR APPLICATION NUMBER: US/09/302,812
 13 <151> PRIOR FILING DATE: 1999-04-30
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 15 <151> PRIOR FILING DATE: 1998-05-01
 16 <160> NUMBER OF SEQ ID NOS: 38
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 19 <211> LENGTH: 4070
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104 35 40 45
105 Phe Arg Val Pro Pro Ser Ser Ser Gly Cys Ala Leu Gly Arg Ala Gly
106 50 55 60
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108 65 70 75 80
109 Thr Ser Trp Met Asp Thr Lys Gly Ile Lys Thr Val Glu Ser Glu Ser
110 85 90 95
111 Leu His Ser Lys Glu Asn Asn Asn Thr Arg Glu Glu Ser Met Met Ser
112 100 105 110
113 Ser Val Gln Lys Asp Asn Phe Tyr Gln His Asn Met Glu Lys Leu Glu
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118 145 150 155 160
119 Glu Gly Pro His Ser Glu Arg Leu Leu Glu Ser Glu Pro Pro Ala Val
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121 Thr Leu Val Pro Glu Gln Phe Ser Asn Ala Asn Val Asp Gln Ser Ser
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123 Pro Lys Asp Asp His Ser Asp Thr Asn Ser Glu Glu Ser Arg Asp Asn
124 195 200 205
125 Gln Gln Phe Leu Thr His Val Lys Leu Ala Asn Ala Lys Gln Thr Met
126 210 215 220
127 Glu Asp Glu Gln Gly Arg Glu Ala Arg Ser His Gln Lys Cys Gly Lys
128 225 230 235 240
129 Ala Cys His Pro Ala Glu Ala Cys Ala Gly Cys Gln Gln Glu Glu Thr
130 245 250 255
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132 260 265 270
133 Gly Thr Gly Leu Lys Asn Ala Asn Arg Leu Asn Arg Gln Glu Ser Ser
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135 Leu Gly Asn Ser Pro Pro Phe Glu Lys Glu Ser Glu Pro Glu Ser Pro
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139 Glu Glu Thr Ser Pro Gly Phe Asp Glu Gln Glu Asp Ser Ser Ala
140 325 330 335
141 Gln Thr Ala Asn Lys Pro Ser Arg Phe Gln Pro Arg Glu Ala Asp Thr
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143 Glu Leu Arg Lys Arg Ser Ser Ala Lys Gly Gly Glu Ile Arg Leu His
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146 370 375 380
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153	His Gln Arg Thr Glu Arg Lys Ile Pro Lys Tyr Ile Pro Pro His Leu			
154	435	440	445	
155	Ser Pro Asp Lys Lys Trp Leu Gly Thr Pro Ile Glu Glu Met Arg Arg			
156	450	455	460	
157	Met Pro Arg Cys Gly Ile Arg Leu Pro Pro Leu Arg Pro Ser Ala Asn			
158	465	470	475	480
159	His Thr Val Thr Ile Arg Val Asp Leu Leu Arg Ile Gly Glu Val Pro			
160	485	490	495	
161	Lys Pro Phe Pro Thr His Phe Lys Asp Leu Trp Asp Asn Lys His Val			
162	500	505	510	
163	Lys Met Pro Cys Ser Glu Gln Asn Leu Tyr Pro Val Glu Asp Glu Asn			
164	515	520	525	
165	Gly Glu Arg Ala Ala Gly Ser Arg Trp Glu Leu Ile Gln Thr Ala Leu			
166	530	535	540	
167	Leu Asn Arg Leu Thr Arg Pro Gln Asn Leu Lys Asp Ala Ile Leu Lys			
168	545	550	555	560
169	Tyr Asn Val Ala Tyr Ser Lys Lys Trp Asp Phe Thr Ala Leu Ile Asp			
170	565	570	575	
171	Phe Trp Asp Lys Val Leu Glu Glu Ala Glu Ala Gln His Leu Tyr Gln			
172	580	585	590	
173	Ser Ile Leu Pro Asp Met Val Lys Ile Ala Leu Cys Leu Pro Asn Ile			
174	595	600	605	
175	Cys Thr Gln Pro Ile Pro Leu Leu Lys Gln Lys Met Asn His Ser Ile			
176	610	615	620	
177	Thr Met Ser Gln Glu Gln Ile Ala Ser Leu Leu Ala Asn Ala Phe Phe			
178	625	630	635	640
179	Cys Thr Phe Pro Arg Arg Asn Ala Lys Met Lys Ser Glu Tyr Ser Ser			
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181	Tyr Pro Asp Ile Asn Phe Asn Arg Leu Phe Glu Gly Arg Ser Ser Arg			
182	660	665	670	
183	Lys Pro Glu Lys Leu Lys Thr Leu Phe Cys Tyr Phe Arg Arg Val Thr			
184	675	680	685	
185	Glu Lys Lys Pro Thr Gly Leu Val Thr Phe Thr Arg Gln Ser Leu Glu			
186	690	695	700	
187	Asp Phe Pro Glu Trp Glu Arg Cys Glu Lys Leu Leu Thr Arg Leu His			
188	705	710	715	720
189	Val Thr Tyr Glu Gly Thr Ile Glu Gly Asn Gly Gln Gly Met Leu Gln			
190	725	730	735	
191	Val Asp Phe Ala Asn Arg Phe Val Gly Gly Gly Val Thr Ser Ala Gly			
192	740	745	750	
193	Leu Val Gln Glu Glu Ile Arg Phe Leu Ile Asn Pro Glu Leu Ile Val			
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195	Ser Arg Leu Phe Thr Glu Val Leu Asp His Asn Glu Cys Leu Ile Ile			
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199 Arg Trp Ala Arg Ser His Glu Asp Arg Ser Glu Arg Asp Asp Trp Gln
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202 820 825 830
203 Tyr Leu Asp Gln Phe Val Pro Glu Lys Ile Arg Arg Glu Leu Asn Lys
204 835 840 845
205 Ala Tyr Cys Gly Phe Leu Arg Pro Gly Val Ser Ser Glu Asn Leu Ser
206 850 855 860
207 Ala Val Ala Thr Gly Asn Trp Gly Cys Gly Ala Phe Gly Gly Asp Ala
208 865 870 875 880
209 Arg Leu Lys Ala Leu Ile Gln Ile Leu Ala Ala Ala Val Ala Glu Arg
210 885 890 895
211 Asp Val Val Tyr Phe Thr Phe Gly Asp Ser Glu Leu Met Arg Asp Ile
212 900 905 910
213 Tyr Ser Met His Thr Phe Leu Thr Glu Arg Lys Leu Thr Val Gly Glu
214 915 920 925
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216 930 935 940
217 Ser Thr Pro Gly Pro Asp Ile Lys Leu Tyr Pro Phe Ile Tyr His Ala
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Use of n and/or Xaa has been detected in the Sequence Listing.
Review the Sequence Listing to insure a corresponding
explanation is presented in the <220> to <223> fields of
each sequence using n or Xaa.

VERIFICATION SUMMARY
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Input Set : N:\Crf3\RULE60\09973451.raw
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L:946 M:258 W: Mandatory Feature missing, <222> not found for SEQ ID#:12
L:946 M:258 W: Mandatory Feature missing, <223> not found for SEQ ID#:12
L:946 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:12
L:964 M:258 W: Mandatory Feature missing, <221> not found for SEQ ID#:14
L:964 M:258 W: Mandatory Feature missing, <222> not found for SEQ ID#:14
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L:964 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:14
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L:1165 M:256 W: Invalid Numeric Header Field, <220> has non-blank data